

CRF Errors Corrected by the STIC System Branch

Serial Number: 08/788,800

CRF Processing Date: 4/02/97
Edited by: WT
Verified by: WT (STIC staff)

Team 3

Changed a file from non-ASCII to ASCII

Changed the margins in cases where the sequence text was "wrapped" down to the next line

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other P0987R1

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as _____

Inserted mandatory headings, specifically: _____

Corrected an obvious error in the response, specifically: _____

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically: _____

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

INPUT SET: S16681.raw

This Raw Listing contains the General Information Section and up to the first 5 pages,

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Bednar, Martin M.
Thomas, G. Roger
Gross, Cordell E.

(ii) TITLE OF INVENTION: ANTI-CD18 ANTIBODIES IN STROKE

(iii) NUMBER OF SEQUENCES: 15

(iv) CORRESPONDENCE ADDRESS:

- (A) ADDRESSEE: Genentech, Inc.
- (B) STREET: 460 Point San Bruno Blvd
- (C) CITY: South San Francisco
- (D) STATE: California
- (E) COUNTRY: USA
- (F) ZIP: 94080

(v) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: WinPatin (Genentech)

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE: 22-Jan-1997
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

- (A) NAME: Lee, Wendy M.
- (B) REGISTRATION NUMBER: 40,378
- (C) REFERENCE/DOCKET NUMBER: P0987rl

(ix) TELECOMMUNICATION INFORMATION:

- (A) TELEPHONE: 415/225-1994
- (B) TELEFAX: 415/952-9881
- (C) TELEX: 910/371-7168

(2) INFORMATION FOR SEO ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 98 amino acids
 - (B) TYPE: Amino Acid

INPUT SET: S16681.raw

47 (D) TOPOLOGY: Linear
48
49 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
50
51 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser
52 1 5 10 15
53
54 Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys
55 20 25 30
56
57 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala
58 35 40 45
59
60 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser
61 50 55 60
62
63 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
64 65 70 75
65
66 Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys Pro Ser
67 80 85 90
68
69 Asn Thr Lys Val Asp Lys Arg Val
70 95 98
71
72 (2) INFORMATION FOR SEQ ID NO:2:
73
74 (i) SEQUENCE CHARACTERISTICS:
75 (A) LENGTH: 98 amino acids
76 (B) TYPE: Amino Acid
77 (D) TOPOLOGY: Linear
78
79 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
80
81 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser
82 1 5 10 15
83
84 Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys Leu Val Lys
85 20 25 30
86
87 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala
88 35 40 45
89
90 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser
91 50 55 60
92
93 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Asn
94 65 70 75
95
96 Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro Ser
97 80 85 90
98
99 Asn Thr Lys Val Asp Lys Thr Val

INPUT SET: S16681.raw

100 95 98
101
102 (2) INFORMATION FOR SEQ ID NO:3:
103
104 (i) SEQUENCE CHARACTERISTICS:
105 (A) LENGTH: 98 amino acids
106 (B) TYPE: Amino Acid
107 (D) TOPOLOGY: Linear
108
109 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
110
111 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser
112 1 5 10 15
113
114 Arg Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys
115 20 25 30
116
117 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala
118 35 40 45
119
120 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser
121 50 55 60
122
123 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
124 65 70 75
125
126 Leu Gly Thr Gln Thr Tyr Thr Cys Asn Val Asn His Lys Pro Ser
127 80 85 90
128
129 Asn Thr Lys Val Asp Lys Arg Val
130 95 98
131
132 (2) INFORMATION FOR SEQ ID NO:4:
133
134 (i) SEQUENCE CHARACTERISTICS:
135 (A) LENGTH: 98 amino acids
136 (B) TYPE: Amino Acid
137 (D) TOPOLOGY: Linear
138
139 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
140
141 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser
142 1 5 10 15
143
144 Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys Leu Val Lys
145 20 25 30
146
147 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala
148 35 40 45
149
150 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser
151 50 55 60
152

INPUT SET: S16681.raw

153 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
154 65 70 75
155
156 Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp His Lys Pro Ser
157 80 85 90
158
159 Asn Thr Lys Val Asp Lys Arg Val
160 95 98
161
162 (2) INFORMATION FOR SEQ ID NO:5:
163
164 (i) SEQUENCE CHARACTERISTICS:
165 (A) LENGTH: 107 amino acids
166 (B) TYPE: Amino Acid
167 (D) TOPOLOGY: Linear
168
169 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
170
171 Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
172 1 5 10 15
173
174 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn
175 20 25 30
176
177 Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn
178 35 40 45
179
180 Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp
181 50 55 60
182
183 Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser
184 65 70 75
185
186 Lys Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr
187 80 85 90
188
189 His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn Arg Gly
190 95 100 105
191
192 Glu Cys
193 107
194
195 (2) INFORMATION FOR SEQ ID NO:6:
196
197 (i) SEQUENCE CHARACTERISTICS:
198 (A) LENGTH: 105 amino acids
199 (B) TYPE: Amino Acid
200 (D) TOPOLOGY: Linear
201
202 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
203
204 Gln Pro Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser
205 1 5 10 15

INPUT SET: S16681.raw

206
207 Glu Glu Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser
208 20 25 30
209
210 Asp Phe Tyr Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser
211 35 40 45
212
213 Ser Pro Val Lys Ala Gly Val Glu Thr Thr Pro Ser Lys Gln
214 50 55 60
215
216 Ser Asn Asn Lys Tyr Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro
217 65 70 75
218
219 Glu Gln Trp Lys Ser His Arg Ser Tyr Ser Cys Gln Val Thr His
220 80 85 90
221
222 Glu Gly Ser Thr Val Glu Lys Thr Val Ala Pro Thr Glu Cys Ser
223 95 100 105
224
225 (2) INFORMATION FOR SEQ ID NO:7:
226
227 (i) SEQUENCE CHARACTERISTICS:
228 (A) LENGTH: 100 amino acids
229 (B) TYPE: Amino Acid
230 (D) TOPOLOGY: Linear
231
232 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
233
234 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Pro
235 1 5 10 15
236
237 Lys Asn Ser Ser Met Ile Ser Asn Thr Pro Ala Leu Gly Cys Leu
238 20 25 30
239
240 Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
241 35 40 45
242
243 Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln
244 50 55 60
245
246 Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro His
247 65 70 75
248
249 Gln Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys
250 80 85 90
251
252 Pro Ser Asn Thr Lys Val Asp Lys Arg Val
253 95 100
254
255 (2) INFORMATION FOR SEQ ID NO:8:
256
257 (i) SEQUENCE CHARACTERISTICS:
258 (A) LENGTH: 11 amino acids

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/788,800

DATE: 04/02/97

TIME: 16:27:50

INPUT SET: S16681.raw

Line

Error

Original Text